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FEB 07 2002

TECH CENTER 1600/2900

SEQUENCE LISTING

<110> DANA-FARBER CANCER INSTITUTE, INC.
MARASCO, Wayne
MHASHILKAR, Abner

<120> INTRABODY-MEDIATED CONTROL OF IMMUNE REACTIONS

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<140> PCT/US98/19563
<141> 1998-09-18

<150> 60/059,339
<151> 1997-09-19

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105/91
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5

10

15

48

gtc ctg tcc cag gtg caa ctg cag cag tca ggg gct gag ctg gca aga
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25

30

96

cct ggg gct tca gtg aag ttg tcc tgc aag gct tct ggc tac acc ttt
Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe

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144

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192

gaa tgg att ggg act att tat cct gga gat ggt gat act agg tac act
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288

aca gcc tac tta cac ctc agc agc ttg tca tct gaa gac tct gcg gtc
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105

110

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INS
g1

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Ala Tyr Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ser Gly Gly Gly	
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Gly Ser Gly Gly Ser Gly Gly Ser Gly Gly Ser Glu Leu Val Leu	
145 150 155 160	
acc caa acc cca acc tcc ctg gct ncc tct ctg gga gac aga gtc acc	528
Thr Gln Thr Pro Thr Ser Leu Ala Xaa Ser Leu Gly Asp Arg Val Thr	
165 170 175	
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Ile Ser Cys Arg Ala Ser Gln Asp Ile Ser Ser Tyr Leu Asn Trp Tyr	
180 185 190	
cag cag aaa cca gat gga act att aaa ctc ctg atc tac tac aca tca	624
Gln Gln Lys Pro Asp Gly Thr Ile Lys Leu Leu Ile Tyr Tyr Thr Ser	
195 200 205	
aga tta tat tca gga gtc cca cca agg ttc agt ggc agt ggg gct gga	672
Arg Leu Tyr Ser Gly Val Pro Pro Arg Phe Ser Gly Ser Gly Ala Gly	
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Thr Asp Tyr Ser Leu Thr Ile Ser Asn Leu Glu Gln Glu Asp Phe Ala	
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Thr Tyr Phe Cys Gln Gln Gly Asn Val Ile Pro Tyr Thr Phe Gly Gly	
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 Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu
 50 55 60
 Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr
 65 70 75 80
 Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr
 85 90 95
 Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val
 100 105 110
 Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe
 115 120 125
 Ala Tyr Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ser Gly Gly Gly
 130 135 140
 Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Glu Leu Val Leu
 145 150 155 160
 Thr Gln Thr Pro Thr Ser Leu Ala Xaa Ser Leu Gly Asp Arg Val Thr
 165 170 175
 Ile Ser Cys Arg Ala Ser Gln Asp Ile Ser Ser Tyr Leu Asn Trp Tyr
 180 185 190
 Gln Gln Lys Pro Asp Gly Thr Ile Lys Leu Leu Ile Tyr Tyr Thr Ser
 195 200 205
 Arg Leu Tyr Ser Gly Val Pro Pro Arg Phe Ser Gly Ser Gly Ala Gly
 210 215 220
 Thr Asp Tyr Ser Leu Thr Ile Ser Asn Leu Glu Gln Glu Asp Phe Ala
 225 230 235 240
 Thr Tyr Phe Cys Gln Gln Gly Asn Val Ile Pro Tyr Thr Phe Gly Gly
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 Glu Lys Asp Glu Leu
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 Val Leu Ser Gln Val Gln Leu Gln Ser Gly Ala Glu Leu Thr Arg
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 Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe
 35 40 45

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 Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu
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gaa tgg att ggg act att tat cct gga gat ggt gat act agg tac act 240
 Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr
 65 70 75 80

cag aat ttc aag ggc aag gcc aca ttg act gca gat aag tcc tcc acc 288
 Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr
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aca gcc tac tta cac ctc agc agc ttg tca tct gaa gac tct gcg gtc 336
 Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val
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 Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe
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 Ala Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Gly Gly Gly
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 Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Leu Gly Asp Thr Ile Thr
 165 170 175

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 Ile Thr Cys His Ala Ser Gln Asn Ile Asn Val Trp Leu Ser Trp Tyr
 180 185 190

cag cag aaa cca gga aat att cct caa cta ttg atc tat aag gct tcc 624
 Gln Gln Lys Pro Gly Asn Ile Pro Gln Leu Leu Ile Tyr Lys Ala Ser
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 245 250 255

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Glu Lys Asp Glu Leu * *
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Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe
35 40 45
Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu
50 55 60
Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr
65 70 75 80
Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr
85 90 95
Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val
100 105
Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe
115 120 125
Ala Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Gly Gly Gly
130 135 140
Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Glu Leu Val Leu
145 150 155 160
Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Leu Gly Asp Thr Ile Thr
165 170 175
Ile Thr Cys His Ala Ser Gln Asn Ile Asn Val Trp Leu Ser Trp Tyr
180 185 190
Gln Gln Lys Pro Gly Asn Ile Pro Gln Leu Leu Ile Tyr Lys Ala Ser
195 200 205
Asn Leu His Thr Gly Val Pro Ser Arg Phe Ser Gly Arg Gly Ser Gly
210 215 220
Thr Gly Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Gly
225 230 235 240
Thr Tyr Tyr Cys Gln Gln Gly Gln Ser Tyr Pro Leu Thr Phe Gly Gly
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Glu Lys Asp Glu Leu

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